

11/15/02

1600

BIOTECHNOLOGY
STEMS
BRANCH

#8

CRF Problem Report

The Scientific and Technical Information Center (STIC) experienced a problem when processing the following computer readable form (CRF):

Application Serial Number: 09/691,344
Filing Date: 10/18/2000
Date Processed by STIC: 2/15/2002

STIC Contact: Mark Spencer, 703-308-4212

Nature of Problem:

The CRF (was):

- ☒ (circle one) Damaged or Unreadable (for Unreadable, see attached)
☐ Blank (no files on CRF) (see attached)
☐ Empty file (filename present, but no bytes in file) (see attached)
☐ Virus-infected. Virus name: _____ The STIC will not process the CRF.
☐ Not saved in ASCII text
☐ Sequence Listing was embedded in the file. According to Sequence Rules, submitted file should **only** be the Sequence Listing.
☐ Did not contain a Sequence Listing. (see attached sample)
☐ Other: _____

**PLEASE USE THE CHECKER VERSION 3.1 PROGRAM TO REDUCE ERRORS.
SEE BELOW FOR ADDRESS:**

<http://www.uspto.gov/web/offices/pac/checker>

→ Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/efc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

R. Mitra

11

5/15/02



1653

RAW SEQUENCE LISTING

DATE: 05/15/2002

PATENT APPLICATION: US/09/691,344A

TIME: 10:59:03

Input Set : N:\CrF3\Refhold\I691344A.raw

Output Set : N:\CRF3\05152002\I691344A.raw

ENTERED

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1 <110> APPLICANT: Donoho, Gregory
2      Turner, C. Alexander Jr.
3      Nehls, Michael
4      Friedrich, Glenn
5      Zambrowicz, Brian
6      Sands, Arthur T.
7 <120> TITLE OF INVENTION: Novel Human Proteins and Polynucleotides
8      Encoding the Same
9 <130> FILE REFERENCE: LEX-0071-USA
C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/691,344A
C--> 11 <141> CURRENT FILING DATE: 2000-10-18
12 <150> PRIOR APPLICATION NUMBER: US 60/160,285
13 <151> PRIOR FILING DATE: 1999-10-19
14 <150> PRIOR APPLICATION NUMBER: US 60/183,583
15 <151> PRIOR FILING DATE: 2000-02-18
16 <160> NUMBER OF SEQ ID NOS: 7
17 <170> SOFTWARE: FastSEQ for Windows Version 4.0
18 <210> SEQ ID NO: 1
19 <211> LENGTH: 1464
20 <212> TYPE: DNA
21 <213> ORGANISM: homo sapiens
22 <400> SEQUENCE: 1
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26 acctgtgctt ctgactatct tctcttcacc agctcttcag atcaatatgg tccatactgt      180
27 ggaagtatga cgtttcccaa agaactcttg ttgaacacaa gtgaagtaac cgtcgcgttt      240
28 gagagtggat cccacatttc tggccggggg tttttgctga cctatgcgag cagcgaccat      300
29 ccagatttaa taacatgttt ggaacgagct agccattatt tgaagacaga atacagcaaa      360
30 ttctgcccgag ctggtttgtg agacgtagca ggagacattt ctgggaatat ggtagatgga      420
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32 ctagggtggc agatcagtggt gcttcagcgc aaagggatca gtcgatatga agggattctg      540
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42 cagattacac aaggtaatga ttcatgtgtg tggcgcaaga caagtcacaa caccagtgtt      1140
43 tcaactaaga aagaagatga gacaatcaca aggcccatcc cctcggaaga aacatccaca      1200
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RAW SEQUENCE LISTING

DATE: 05/15/2002

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TIME: 10:59:03

Input Set : N:\CRf3\Refhold\I691344A.raw

Output Set: N:\CRF3\05152002\I691344A.raw

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46      gcggaggctc agaaaacaga ctgttggaag cagattaaat atccctttgc cagacatcag      1380
47      tcagctgagt ttaccatcag ctatgataat gagaaggaga tgacacaaaa gttagatctc      1440
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53 <213> ORGANISM: homo sapiens
54 <400> SEQUENCE: 2
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57      Glu Lys Thr Ile Thr Val Pro Lys Gly Lys Arg Leu Ile Leu Arg Leu
58      20              25              30
59      Gly Asp Leu Asp Ile Glu Ser Gln Thr Cys Ala Ser Asp Tyr Leu Leu
60      35              40              45
61      Phe Thr Ser Ser Ser Asp Gln Tyr Gly Pro Tyr Cys Gly Ser Met Thr
62      50              55              60
63      Val Pro Lys Glu Leu Leu Asn Thr Ser Glu Val Thr Val Arg Phe
64      65              70              75              80
65      Glu Ser Gly Ser His Ile Ser Gly Arg Gly Phe Leu Leu Thr Tyr Ala
66      85              90              95
67      Ser Ser Asp His Pro Asp Leu Ile Thr Cys Leu Glu Arg Ala Ser His
68      100             105             110
69      Tyr Leu Lys Thr Glu Tyr Ser Lys Phe Cys Pro Ala Gly Cys Arg Asp
70      115             120             125
71      Val Ala Gly Asp Ile Ser Gly Asn Met Val Asp Gly Tyr Arg Asp Thr
72      130             135             140
73      Ser Leu Leu Cys Lys Ala Ala Ile His Ala Gly Ile Ile Ala Asp Glu
74      145             150             155             160
75      Leu Gly Gly Gln Ile Ser Val Leu Gln Arg Lys Gly Ile Ser Arg Tyr
76      165             170             175
77      Glu Gly Ile Leu Ala Asn Gly Val Leu Ser Arg Asp Gly Ser Leu Ser
78      180             185             190
79      Asp Lys Arg Phe Leu Phe Thr Ser Asn Gly Cys Ser Arg Ser Leu Ser
80      195             200             205
81      Phe Glu Pro Asp Gly Gln Ile Arg Ala Ser Ser Ser Trp Gln Ser Val
82      210             215             220
83      Asn Glu Ser Gly Asp Gln Val His Trp Ser Pro Gly Gln Ala Arg Leu
84      225             230             235             240
85      Gln Asp Gln Gly Pro Ser Trp Ala Ser Gly Asp Ser Ser Asn Asn His
86      245             250             255
87      Lys Pro Arg Glu Trp Leu Glu Ile Asp Leu Gly Glu Lys Lys Lys Ile
88      260             265             270
89      Thr Gly Ile Arg Thr Thr Gly Ser Thr Gln Ser Asn Phe Asn Phe Tyr
90      275             280             285
91      Val Lys Ser Phe Val Met Asn Phe Lys Asn Asn Asn Ser Lys Trp Lys
92      290             295             300
93      Thr Tyr Lys Gly Ile Val Asn Asn Glu Glu Lys Val Phe Gln Gly Asn
94      305             310             315             320

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RAW SEQUENCE LISTING

DATE: 05/15/2002

PATENT APPLICATION: US/09/691,344A

TIME: 10:59:03

Input Set : N:\Crf3\Refhold\I691344A.raw

Output Set: N:\CRF3\05152002\I691344A.raw

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95      Ser Asn Phe Arg Asp Pro Val Gln Asn Asn Phe Ile Pro Pro Ile Val
96              325              330              335
97      Ala Arg Tyr Val Arg Val Val Pro Gln Thr Trp His Gln Arg Ile Ala
98              340              345              350
99      Leu Lys Val Glu Leu Ile Gly Cys Gln Ile Thr Gln Gly Asn Asp Ser
100             355              360              365
101      Leu Val Trp Arg Lys Thr Ser Gln Ser Thr Ser Val Ser Thr Lys Lys
102             370              375              380
103      Glu Asp Glu Thr Ile Thr Arg Pro Ile Pro Ser Glu Glu Thr Ser Thr
104             385              390              395              400
105      Gly Ile Asn Ile Thr Thr Val Ala Ile Pro Leu Val Leu Val Val
106             405              410              415
107      Leu Val Phe Ala Gly Met Gly Ile Phe Ala Ala Phe Arg Lys Lys Lys
108             420              425              430
109      Lys Lys Gly Ser Pro Tyr Gly Ser Ala Glu Ala Gln Lys Thr Asp Cys
110             435              440              445
111      Trp Lys Gln Ile Lys Tyr Pro Phe Ala Arg His Gln Ser Ala Glu Phe
112             450              455              460
113      Thr Ile Ser Tyr Asp Asn Glu Lys Glu Met Thr Gln Lys Leu Asp Leu
114             465              470              475              480
115      Ile Thr Ser Asp Met Ala Gly
116             485
118 <210> SEQ ID NO: 3
119 <211> LENGTH: 1761
120 <212> TYPE: DNA
121 <213> ORGANISM: homo sapiens
122 <400> SEQUENCE: 3
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124      gaggcggccc ggcccgggca gctgcggctc gggatccgtc gaggggaggc cgagcttgcc      120
125      aagctggcgc ccacggggt catggtgcc cgcgcgcgcg cgggcggcgc actggcgcg      180
126      gctgcggggc ggggcctct ggctttgctg ctgcgggtct ccgcccgct ccggtgcag      240
127      gcggaggagc tgggtgatgg ctgtggacac ctagtgaact atcaggatag tggcacaatg      300
128      acatctaaga attatcccgc gacctacccc aatcacactg ttgcgaaaa gacaattaca      360
129      gtaccaaagg gaaaaagact gattctgagg ttgggagatt tggatatcga atcccagacc      420
130      tgtgcttctg actatcttct cttcaccagc tcttcagatc aatatggtcc atactgtgga      480
131      agtatgactg ttcccaaaga actcttggtg aacacaagtg aagtaaccgt ccgctttgag      540
132      agtggatccc acatttcttg ccggggtttt ttgctgacct atgcgagcag cgaccatcca      600
133      gatttaataa catgtttgga acgagctagc cattatttga agacagaata cagcaaattc      660
134      tgccagctcg gttgtagaga cgtagcagga gacatttctg ggaatatggt agatggatat      720
135      agagatacct ctttattgtg caaagctgcc atccatgcag gaataattgc tgatgaacta      780
136      ggtggccaga tcagtggtgt tcacgcgcaa gggatcagtc gatatgaagg gattctggcc      840
137      aatggtgttc tttcgaggga tggttccctg tcagacaagc gatttctggt taccctccat      900
138      ggttcagca gatccttgag ttttgaacct gacgggcaaa tcagagcttc ttcctcatgg      960
139      cagtcggtca atgagagtgg agaccaagtt cactggtctc ctggccaagc ccgacttcag      1020
140      gaccaaggc catcatgggc ttccggcgac agtagcaaca accacaaacc acgagagtgg      1080
141      ctggagatcg atttggggga gaaaaagaaa ataacaggaa ttaggaccac aggatctaca      1140
142      cagtcgaact tcaactttta tgttaagagt ttgtgatga acttcaaaaa caataattct      1200
143      aagtgaaga cctataaagg aattgtgaat aatgaagaaa aggtgtttca gggtaaactct      1260
144      aactttcggg acccagtgca aaacaatttc atccctccca tcgtggccag atatgtgcgg      1320

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RAW SEQUENCE LISTING

DATE: 05/15/2002

PATENT APPLICATION: US/09/691,344A

TIME: 10:59:04

Input Set : N:\Crf3\Refhold\I691344A.raw

Output Set: N:\CRF3\05152002\I691344A.raw

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147      actaagaaag aagatgagac aatcacaagg cccatcccct cggaagaaac atccacagga      1500
148      ataaacatta caacggtggc tattccattg gtgtcccttg ttgtcctggt gtttgctgga      1560
149      atggggatct ttgcagcctt tagaaagaag aagaagaaag gaagtccgta tggatcagcg      1620
150      gaggctcaga aaacagactg ttggaagcag attaaatatc cctttgccag acatcagtca      1680
151      gctgagttta ccatcagcta tgataatgag aaggagatga cacaaaagtt agatctcatc      1740
152      acaagtgata tggcaggtta a                                     1761
154 <210> SEQ ID NO: 4
155 <211> LENGTH: 586
156 <212> TYPE: PRT
157 <213> ORGANISM: homo sapiens
158 <400> SEQUENCE: 4
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160      1          5          10          15
161      Ser Ser Ala Glu Glu Ala Ala Arg Pro Gly Gln Leu Arg Leu Gly Ile
162      20          25          30
163      Arg Arg Gly Glu Ala Glu Leu Ala Lys Leu Ala Pro Ser Gly Val Met
164      35          40          45
165      Val Pro Gly Ala Arg Gly Gly Ala Leu Ala Arg Ala Ala Gly Arg
166      50          55          60
167      Gly Leu Leu Ala Leu Leu Leu Ala Val Ser Ala Pro Leu Arg Leu Gln
168      65          70          75          80
169      Ala Glu Glu Leu Gly Asp Gly Cys Gly His Leu Val Thr Tyr Gln Asp
170      85          90          95
171      Ser Gly Thr Met Thr Ser Lys Asn Tyr Pro Gly Thr Tyr Pro Asn His
172      100         105         110
173      Thr Val Cys Glu Lys Thr Ile Thr Val Pro Lys Gly Lys Arg Leu Ile
174      115         120         125
175      Leu Arg Leu Gly Asp Leu Asp Ile Glu Ser Gln Thr Cys Ala Ser Asp
176      130         135         140
177      Tyr Leu Leu Phe Thr Ser Ser Ser Asp Gln Tyr Gly Pro Tyr Cys Gly
178      145         150         155         160
179      Ser Met Thr Val Pro Lys Glu Leu Leu Leu Asn Thr Ser Glu Val Thr
180      165         170         175
181      Val Arg Phe Glu Ser Gly Ser His Ile Ser Gly Arg Gly Phe Leu Leu
182      180         185         190
183      Thr Tyr Ala Ser Ser Asp His Pro Asp Leu Ile Thr Cys Leu Glu Arg
184      195         200         205
185      Ala Ser His Tyr Leu Lys Thr Glu Tyr Ser Lys Phe Cys Pro Ala Gly
186      210         215         220
187      Cys Arg Asp Val Ala Gly Asp Ile Ser Gly Asn Met Val Asp Gly Tyr
188      225         230         235         240
189      Arg Asp Thr Ser Leu Cys Lys Ala Ala Ile His Ala Gly Ile Ile
190      245         250         255
191      Ala Asp Glu Leu Gly Gly Gln Ile Ser Val Leu Gln Arg Lys Gly Ile
192      260         265         270
193      Ser Arg Tyr Glu Gly Ile Leu Ala Asn Gly Val Leu Ser Arg Asp Gly
194      275         280         285

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RAW SEQUENCE LISTING

DATE: 05/15/2002

PATENT APPLICATION: US/09/691,344A

TIME: 10:59:04

Input Set : N:\Crf3\Refhold\I691344A.raw

Output Set: N:\CRF3\05152002\I691344A.raw

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195      Ser Leu Ser Asp Lys Arg Phe Leu Phe Thr Ser Asn Gly Cys Ser Arg
196          290                      295                      300
197      Ser Leu Ser Phe Glu Pro Asp Gly Gln Ile Arg Ala Ser Ser Ser Trp
198          305                      310                      315                      320
199      Gln Ser Val Asn Glu Ser Gly Asp Gln Val His Trp Ser Pro Gly Gln
200                      325                      330                      335
201      Ala Arg Leu Gln Asp Gln Gly Pro Ser Trp Ala Ser Gly Asp Ser Ser
202                      340                      345                      350
203      Asn Asn His Lys Pro Arg Glu Trp Leu Glu Ile Asp Leu Gly Glu Lys
204                      355                      360                      365
205      Lys Lys Ile Thr Gly Ile Arg Thr Thr Gly Ser Thr Gln Ser Asn Phe
206          370                      375                      380
207      Asn Phe Tyr Val Lys Ser Phe Val Met Asn Phe Lys Asn Asn Asn Ser
208          385                      390                      395                      400
209      Lys Trp Lys Thr Tyr Lys Gly Ile Val Asn Asn Glu Glu Lys Val Phe
210                      405                      410                      415
211      Gln Gly Asn Ser Asn Phe Arg Asp Pro Val Gln Asn Asn Phe Ile Pro
212                      420                      425                      430
213      Pro Ile Val Ala Arg Tyr Val Arg Val Val Pro Gln Thr Trp His Gln
214          435                      440                      445
215      Arg Ile Ala Leu Lys Val Glu Leu Ile Gly Cys Gln Ile Thr Gln Gly
216          450                      455                      460
217      Asn Asp Ser Leu Val Trp Arg Lys Thr Ser Gln Ser Thr Ser Val Ser
218          465                      470                      475                      480
219      Thr Lys Lys Glu Asp Glu Thr Ile Thr Arg Pro Ile Pro Ser Glu Glu
220                      485                      490                      495
221      Thr Ser Thr Gly Ile Asn Ile Thr Thr Val Ala Ile Pro Leu Val Leu
222          500                      505                      510
223      Leu Val Val Leu Val Phe Ala Gly Met Gly Ile Phe Ala Ala Phe Arg
224          515                      520                      525
225      Lys Lys Lys Lys Lys Gly Ser Pro Tyr Gly Ser Ala Glu Ala Gln Lys
226          530                      535                      540
227      Thr Asp Cys Trp Lys Gln Ile Lys Tyr Pro Phe Ala Arg His Gln Ser
228          545                      550                      555                      560
229      Ala Glu Phe Thr Ile Ser Tyr Asp Asn Glu Lys Glu Met Thr Gln Lys
230                      565                      570                      575
231      Leu Asp Leu Ile Thr Ser Asp Met Ala Gly
232          580                      585
233
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236 <212> TYPE: DNA
237 <213> ORGANISM: homo sapiens
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242      acctacccca atcacactgt ttgcgaaaag acaattacag taccaaaagg gaaaagactg      240
243      attctgaggt tgggagattt ggatatcgaa tccagacct gtgcttctga ctatcttctc      300
244      ttcaccagct cttcagatca atatggtcca tactgtggaa gtatgactgt tcccaaagaa      360

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VERIFICATION SUMMARY

DATE: 05/15/2002

PATENT APPLICATION: US/09/691,344A

TIME: 10:59:05

Input Set : N:\Crf3\Refhold\I691344A.raw

Output Set: N:\CRF3\05152002\I691344A.raw

L:10 M:270 C: Current Application Number differs, Wrong Format

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date